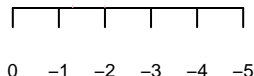
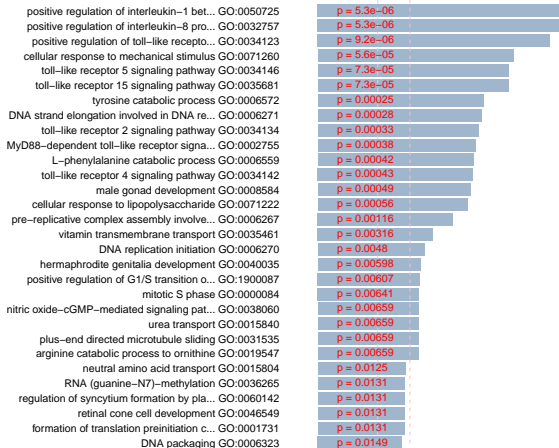


Oecy.Fem-Mal_difpos top GO: BP fisher elin

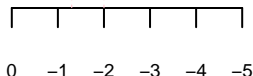


log(p)

n=46/72

Oecy.Fem-Mal_difpos top GO: MF fisher elin

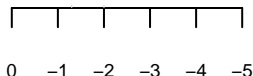
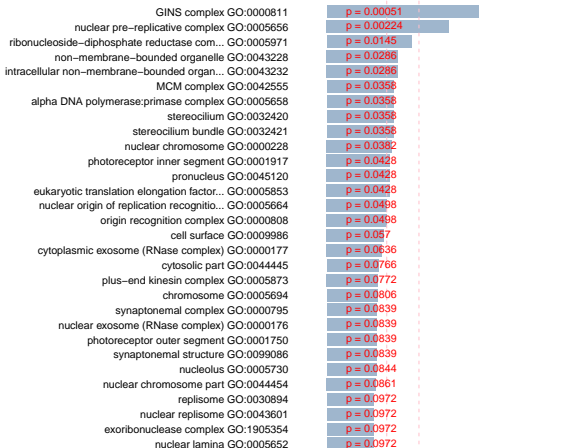
sodium-dependent multivitamin transmembr...	GO:0008523	p = 0.0015
tRNA (guanine-N7-)-methyltransferase act...	GO:0008176	p = 0.0066
glycine C-acetyltransferase activity	GO:0008890	p = 0.0066
tyrosine 3-monooxygenase activity	GO:0004511	p = 0.0066
TBP-class protein binding	GO:0017025	p = 0.0066
homogentisate 1,2-dioxygenase activity	GO:0004411	p = 0.0066
4-hydroxyphenylpyruvate dioxygenase acti...	GO:0003868	p = 0.0066
beta-glucuronidase activity	GO:0004566	p = 0.0066
dUTP diphosphatase activity	GO:0004170	p = 0.0066
ribonucleoside-diphosphate reductase act...	GO:0061731	p = 0.0132
DNA ligase activity	GO:0003909	p = 0.0132
ribonucleoside-diphosphate reductase act...	GO:0004748	p = 0.0132
proline dehydrogenase activity	GO:0004657	p = 0.0132
myostimulatory hormone activity	GO:0016084	p = 0.0132
oxidoreductase activity, acting on CH or...	GO:0016728	p = 0.0132
DNA ligase (ATP) activity	GO:0003910	p = 0.0132
arginase activity	GO:0004053	p = 0.0132
diuretic hormone activity	GO:0008613	p = 0.0198
eukaryotic initiation factor 4E binding	GO:0008190	p = 0.0198
structural constituent of adult chitin-b...	GO:0008012	p = 0.0198
neuropeptide receptor binding	GO:0071855	p = 0.0198
hydrolase activity, acting on carbon-nit...	GO:0016813	p = 0.0263
SAM domain binding	GO:0032093	p = 0.0263
pyridoxal phosphate binding	GO:0030170	p = 0.0263
vitamin B6 binding	GO:0070279	p = 0.0263
ligase activity, forming phosphoric este...	GO:0016886	p = 0.0263
UDP-glucose 4-epimerase activity	GO:0003978	p = 0.0328
racemase and epimerase activity, acting ...	GO:0016857	p = 0.0328
calcium:sodium antiporter activity	GO:0005432	p = 0.0328
structural molecule activity	GO:0005198	p = 0.0392



log(p)

n=39/72

Oecy.Fem-Mal_difpos top GO: CC fisher elin



log(p)

n=43/72